

09/807452
PCT/US99/24511

WO 00/23589

SEQUENCE LISTING

<110> INCYTE PHARMACEUTICALS, INC.

TANG, Y. Tom
YUE, Henry
HILLMAN, Jennifer L.
GUEGLER, Karl J.
CORLEY, Neil C.
LAL, Preeti
AZIMZAI, Yalda
BAUGHN, Mariah R.
JUNMING, Yang
SHIH, Leo L.

<120> PROLIFERATION AND APOPTOSIS RELATED PROTEINS

<130> PF-0619 PCT

<140> To Be Assigned

<141> Herewith

<150> 09/175,737; unassigned; 60/118,559; 09/249,740; unassigned;
60/154,336<151> 1998-10-20; 1998-10-20; 1999-02-04; 1999-04-11; 1999-04-11;
1999-04-22

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<170> PERL Program

<210> 1

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<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 1342011CD1

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Val Thr Gly Ala Gly Gly Trp Gly Ser Ala Ala Val Cys Arg Gly
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Arg Ala Leu Arg Gly Arg Glu Pro Ala Leu Pro Ser Ala Ser Phe
35 40 45
Pro Asp Val Ala Ala Cys Pro Gly Ser Leu Asp Cys Ala Leu Lys
50 55 60
Arg Arg Ala Arg Cys Pro Pro Gly Ala His Ala Cys Gly Pro Cys
65 70 75
Leu Gln Pro Phe Gln Glu Asp Gln Gln Gly Leu Cys Val Pro Arg
80 85 90
Met Arg Arg Pro Pro Gly Gly Arg Pro Gln Pro Arg Leu Glu
95 100 105

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Asp	Glu	Ile	Asp	Phe	Leu	Ala	Gln	Glu	Leu	Ala	Arg	Lys	Glu	Ser
				110				115					120	
Gly	His	Ser	Thr	Pro	Pro	Leu	Pro	Lys	Asp	Arg	Gln	Arg	Leu	Pro
				125				130					135	
Glu	Pro	Ala	Thr	Leu	Gly	Phe	Ser	Ala	Arg	Gly	Gln	Gly	Leu	Glu
				140				145					150	
Leu	Gly	Leu	Pro	Ser	Thr	Pro	Gly	Thr	Pro	Thr	Pro	Thr	Pro	His
				155				160					165	
Thr	Ser	Leu	Gly	Ser	Pro	Val	Ser	Ser	Asp	Pro	Val	His	Met	Ser
				170				175					180	
Pro	Leu	Glu	Pro	Arg	Gly	Gly	Gln	Gly	Asp	Gly	Leu	Ala	Leu	Val
				185				190					195	
Leu	Ile	Leu	Ala	Phe	Cys	Val	Ala	Gly	Ala	Ala	Ala	Leu	Ser	Val
				200				205					210	
Ala	Ser	Leu	Cys	Trp	Cys	Arg	Leu	Gln	Arg	Glu	Ile	Arg	Leu	Thr
				215				220					225	
Gln	Lys	Ala	Asp	Tyr	Ala	Thr	Ala	Lys	Ala	Pro	Gly	Ser	Pro	Ala
				230				235					240	
Ala	Pro	Arg	Ile	Ser	Pro	Gly	Asp	Gln	Arg	Leu	Ala	Gln	Ser	Ala
				245				250					255	
Glu	Met	Tyr	His	Tyr	Gln	His	Gln	Arg	Gln	Gln	Met	Leu	Cys	Leu
				260				265					270	
Glu	Arg	His	Lys	Glu	Pro	Pro	Lys	Glu	Leu	Asp	Thr	Ala	Ser	Ser
				275				280					285	
Asp	Glu	Glu	Asn	Glu	Asp	Gly	Asp	Phe	Thr	Val	Tyr	Glu	Cys	Pro
				290				295					300	
Gly	Leu	Ala	Pro	Thr	Gly	Glu	Met	Glu	Val	Arg	Asn	Pro	Leu	Phe
				305				310					315	
Asp	His	Ala	Ala	Leu	Ser	Ala	Pro	Leu	Pro	Ala	Pro	Ser	Ser	Pro
				320				325					330	
Pro	Ala	Leu	Pro											

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<220>
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Ser Arg His Asp Met Leu Ala Trp Val Asn Asp Ser Leu His Leu
20 25 30
Asn Tyr Thr Lys Ile Glu Gln Leu Cys Ser Gly Ala Ala Tyr Cys
35 40 45
Gln Phe Met Asp Met Leu Phe Pro Gly Cys Val His Leu Arg Lys
50 55 60
Val Lys Phe Gln Ala Lys Leu Glu His Glu Tyr Ile His Asn Phe
65 70 75
Lys Val Leu Gln Ala Ala Phe Lys Lys Met Gly Val Asp Lys Ile
80 85 90

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Ile	Pro	Val	Glu	Lys	Leu	Val	Lys	Gly	Lys	Phe	Gln	Asp	Asn	Phe
			95				100							105
Glu	Phe	Ile	Gln	Trp	Phe	Lys	Lys	Phe	Phe	Asp	Ala	Asn	Tyr	Asp
			110				115							120
Gly	Lys	Asp	Tyr	Asn	Pro	Leu	Leu	Ala	Arg	Gln	Gly	Gln	Asp	Val
			125				130							135
Ala	Pro	Pro	Pro	Asn	Pro	Gly	Asp	Gln	Ile	Phe	Asn	Lys	Ser	Lys
			140				145							150
Lys	Leu	Ile	Gly	Thr	Ala	Val	Pro	Gln	Arg	Thr	Ser	Pro	Thr	Gly
			155				160							165
Pro	Lys	Asn	Met	Gln	Thr	Ser	Gly	Arg	Leu	Ser	Asn	Val	Ala	Pro
			170				175							180
Pro	Cys	Ile	Leu	Arg	Lys	Asn	Pro	Pro	Ser	Ala	Arg	Asn	Gly	Gly
			185				190							195
His	Glu	Thr	Asp	Ala	Gln	Ile	Leu	Glu	Leu	Asn	Gln	Gln	Leu	Val
			200				205							210
Asp	Leu	Lys	Leu	Thr	Val	Asp	Gly	Leu	Glu	Lys	Glu	Arg	Asp	Phe
			215				220							225
Tyr	Phe	Ser	Lys	Leu	Arg	Asp	Ile	Glu	Leu	Ile	Cys	Gln	Glu	His
			230				235							240
Glu	Ser	Glu	Asn	Ser	Pro	Val	Ile	Ser	Gly	Ile	Ile	Gly	Ile	Leu
			245				250							255
Tyr	Ala	Thr	Glu	Glu	Gly	Phe	Ala	Pro	Pro	Glu	Asp	Asp	Glu	Ile
			260				265							270
Glu	Glu	His	Gln	Gln	Glu	Asp	Gln	Asp	Glu	Tyr				
			275				280							

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Ser Asp Met Arg Gln Glu Lys Pro Ser Ser Pro Ser Pro Met Pro
20 25 30
Ser Ser Thr Pro Ser Pro Ser Leu Asn Leu Gly Asn Thr Glu Glu
35 40 45
Ala Ile Arg Asp Asn Ser Gln Val Asn Ala Val Thr Val Leu Thr
50 55 60
Leu Leu Asp Lys Leu Val Asn Met Leu Asp Ala Val Gln Glu Asn
65 70 75
Gln His Lys Met Glu Gln Arg Gln Ile Ser Leu Glu Gly Ser Val
80 85 90
Lys Gly Ile Gln Asn Asp Leu Thr Lys Leu Ser Lys Tyr Gln Ala
95 100 105
Ser Thr Ser Asn Thr Val Ser Lys Leu Leu Glu Lys Ser Arg Lys
110 115 120
Val Ser Ala His Thr Arg Ala Val Lys Glu Arg Met Asp Arg Gln

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125	130	135
Cys Ala Gln Val	Lys Arg Leu Glu Asn	Asn His Ala Gln Leu
140	145	150
Arg Arg Asn His	Phe Lys Val Leu Ile	Phe Gln Glu Glu Asn Glu
155	160	165
Ile Pro Ala Ser	Val Phe Val Lys Gln	Pro Val Ser Gly Ala Val
170	175	180
Glu Gly Lys Glu	Glu Leu Pro Asp Glu	Asn Lys Ser Leu Glu Glu
185	190	195
Thr Leu His Thr	Val Asp Leu Ser Ser	Asp Asp Asp Leu Pro His
200	205	210
Asp Glu Glu Ala	Leu Glu Asp Ser Ala	Glu Glu Lys Val Gly Arg
215	220	225
Ser Pro Arg Gly	Arg Glu Ile Lys Arg	Ser Arg Pro
230	235	

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<211> 941

<212> PRT

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<220>

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<223> Incyte ID No: 939000CD1

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Met Asn Lys Lys Lys	Pro Phe Leu Gly	Met Pro Ala Pro Leu
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Gly Tyr Val Pro Gly	Leu Gly Arg Gly	Ala Thr Gly Phe Thr Thr
20	25	30
Arg Ser Asp Ile Gly	Pro Ala Arg Asp	Ala Asn Asp Pro Val Asp
35	40	45
Asp Arg His Ala Pro	Pro Gly Lys Arg	Thr Val Gly Asp Gln Met
50	55	60
Lys Lys Asn Gln Ala	Ala Asp Asp Asp	Asp Glu Asp Leu Asn Asp
65	70	75
Thr Asn Tyr Asp Glu	Phe Asn Gly Tyr	Ala Gly Ser Leu Phe Ser
80	85	90
Ser Gly Pro Tyr Glu	Lys Asp Asp Glu	Glu Ala Asp Ala Ile Tyr
95	100	105
Ala Ala Leu Asp Lys	Arg Met Asp Glu	Arg Arg Lys Glu Arg Arg
110	115	120
Glu Gln Arg Glu Lys	Glu Glu Ile Glu	Lys Tyr Arg Met Glu Arg
125	130	135
Pro Lys Ile Gln Gln	Gln Phe Ser Asp	Leu Lys Arg Lys Leu Ala
140	145	150
Glu Val Thr Glu Glu	Trp Leu Ser Ile	Pro Glu Val Gly Asp
155	160	165
Ala Arg Asn Lys Arg	Gln Arg Asn Pro	Arg Tyr Glu Lys Leu Thr
170	175	180
Pro Val Pro Asp Ser	Phe Phe Ala Lys	His Leu Gln Thr Gly Glu
185	190	195
Asn His Thr Ser Val	Asp Pro Arg Gln	Thr Gln Phe Gly Gly Leu
200	205	210

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Asn	Thr	Pro	Tyr	Pro	Gly	Gly	Leu	Asn	Thr	Pro	Tyr	Pro	Gly	Gly
					215				220					225
Met	Thr	Pro	Gly	Leu	Met	Thr	Pro	Gly	Thr	Gly	Glu	Leu	Asp	Met
					230				235					240
Arg	Lys	Ile	Gly	Gln	Ala	Arg	Asn	Thr	Leu	Met	Asp	Met	Arg	Leu
					245				250					255
Ser	Gln	Val	Ser	Asp	Ser	Val	Ser	Gly	Gln	Thr	Val	Val	Asp	Pro
					260				265					270
Lys	Gly	Tyr	Leu	Thr	Asp	Leu	Asn	Ser	Met	Ile	Pro	Thr	His	Gly
					275				280					285
Gly	Asp	Ile	Asn	Asp	Ile	Lys	Lys	Ala	Arg	Leu	Leu	Leu	Lys	Ser
					290				295					300
Val	Arg	Glu	Thr	Asn	Pro	His	His	Pro	Pro	Ala	Trp	Ile	Ala	Ser
					305				310					315
Ala	Arg	Leu	Glu	Glu	Val	Thr	Gly	Lys	Leu	Gln	Val	Ala	Arg	Asn
					320				325					330
Leu	Ile	Met	Lys	Gly	Thr	Glu	Met	Cys	Pro	Lys	Ser	Glu	Asp	Val
					335				340					345
Trp	Leu	Glu	Ala	Ala	Arg	Leu	Gln	Pro	Gly	Asp	Thr	Ala	Lys	Ala
					350				355					360
Val	Val	Ala	Gln	Ala	Val	Arg	His	Leu	Pro	Gln	Ser	Val	Arg	Ile
					365				370					375
Tyr	Ile	Arg	Ala	Ala	Glu	Leu	Glu	Thr	Asp	Ile	Arg	Ala	Lys	Lys
					380				385					390
Arg	Val	Leu	Arg	Lys	Ala	Leu	Glu	His	Val	Pro	Asn	Ser	Val	Arg
					395				400					405
Leu	Trp	Lys	Ala	Ala	Val	Glu	Leu	Glu	Glu	Pro	Glu	Asp	Ala	Arg
					410				415					420
Ile	Met	Leu	Ser	Arg	Ala	Val	Glu	Cys	Cys	Pro	Thr	Ser	Val	Glu
					425				430					435
Leu	Trp	Leu	Ala	Leu	Ala	Arg	Leu	Glu	Thr	Tyr	Glu	Asn	Ala	Arg
					440				445					450
Lys	Val	Leu	Asn	Lys	Ala	Arg	Glu	Asn	Ile	Pro	Thr	Asp	Arg	His
					455				460					465
Ile	Trp	Ile	Thr	Ala	Ala	Lys	Leu	Glu	Glu	Ala	Asn	Gly	Asn	Thr
					470				475					480
Gln	Met	Val	Glu	Lys	Ile	Ile	Asp	Arg	Ala	Ile	Thr	Ser	Leu	Arg
					485				490					495
Ala	Asn	Gly	Val	Glu	Ile	Asn	Arg	Glu	Gln	Trp	Ile	Gln	Asp	Ala
					500				505					510
Glu	Glu	Cys	Asp	Arg	Ala	Gly	Ser	Val	Ala	Thr	Cys	Gln	Ala	Val
					515				520					525
Met	Arg	Ala	Val	Ile	Gly	Ile	Gly	Ile	Glu	Glu	Glu	Asp	Arg	Lys
					530				535					540
His	Thr	Trp	Met	Glu	Asp	Ala	Asp	Ser	Cys	Val	Ala	His	Asn	Ala
					545				550					555
Leu	Glu	Cys	Ala	Arg	Ala	Ile	Tyr	Ala	Tyr	Ala	Leu	Gln	Val	Phe
					560				565					570
Pro	Ser	Lys	Lys	Ser	Val	Trp	Leu	Arg	Ala	Ala	Tyr	Phe	Glu	Lys
					575				580					585
Asn	His	Gly	Thr	Arg	Glu	Ser	Leu	Glu	Ala	Leu	Leu	Gln	Arg	Ala
					590				595					600
Val	Ala	His	Cys	Pro	Lys	Ala	Glu	Val	Leu	Trp	Leu	Met	Gly	Ala
					605				610					615
Lys	Ser	Lys	Trp	Leu	Ala	Gly	Asp	Val	Pro	Ala	Ala	Arg	Ser	Ile

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	620	625	630											
Leu	Ala	Leu	Ala	Phe	Gln	Ala	Asn	Pro	Asn	Ser	Glu	Glu	Ile	Trp
				635		640								645
Leu	Ala	Ala	Val	Lys	Leu	Glu	Ser	Glu	Asn	Asp	Glu	Tyr	Glu	Arg
				650		655								660
Ala	Arg	Arg	Leu	Leu	Ala	Lys	Ala	Arg	Ser	Ser	Ala	Pro	Thr	Ala
				665		670								675
Arg	Val	Phe	Met	Lys	Ser	Val	Lys	Leu	Glu	Trp	Val	Gln	Asp	Asn
				680		685								690
Ile	Arg	Ala	Ala	Gln	Asp	Leu	Cys	Glu	Glu	Ala	Leu	Arg	His	Tyr
				695		700								705
Glu	Asp	Phe	Pro	Lys	Leu	Trp	Met	Met	Lys	Gly	Gln	Ile	Glu	Glu
				710		715								720
Gln	Lys	Glu	Met	Met	Glu	Lys	Ala	Arg	Glu	Ala	Tyr	Asn	Gln	Gly
				725		730								735
Leu	Lys	Lys	Cys	Pro	His	Ser	Thr	Pro	Leu	Trp	Leu	Leu	Leu	Ser
				740		745								750
Arg	Leu	Glu	Glu	Lys	Ile	Gly	Gln	Leu	Thr	Arg	Ala	Arg	Ala	Ile
				755		760								765
Leu	Glu	Lys	Ser	Arg	Leu	Lys	Asn	Pro	Lys	Asn	Pro	Gly	Leu	Trp
				770		775								780
Leu	Glu	Ser	Val	Arg	Leu	Glu	Tyr	Arg	Ala	Gly	Leu	Lys	Asn	Ile
				785		790								795
Ala	Asn	Thr	Leu	Met	Ala	Lys	Ala	Leu	Gln	Glu	Cys	Pro	Asn	Ser
				800		805								810
Gly	Ile	Leu	Trp	Ser	Glu	Ala	Ile	Phe	Leu	Glu	Ala	Arg	Pro	Gln
				815		820								825
Arg	Arg	Thr	Lys	Ser	Val	Asp	Ala	Leu	Lys	Lys	Cys	Glu	His	Asp
				830		835								840
Pro	His	Val	Leu	Leu	Ala	Val	Ala	Lys	Leu	Phe	Trp	Ser	Gln	Arg
				845		850								855
Lys	Ile	Thr	Lys	Ala	Arg	Glu	Trp	Phe	His	Arg	Thr	Val	Lys	Ile
				860		865								870
Asp	Ser	Asp	Leu	Gly	Asp	Ala	Trp	Ala	Phe	Phe	Tyr	Lys	Phe	Glu
				875		880								885
Leu	Gln	His	Gly	Thr	Glu	Glu	Gln	Gln	Glu	Glu	Val	Arg	Lys	Arg
				890		895								900
Cys	Glu	Ser	Ala	Glu	Pro	Arg	His	Gly	Glu	Leu	Trp	Cys	Ala	Val
				905		910								915
Ser	Lys	Asp	Ile	Ala	Asn	Trp	Gln	Lys	Lys	Ile	Gly	Asp	Ile	Leu
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Arg	Leu	Val	Ala	Gly	Arg	Ile	Lys	Asn	Thr	Phe				
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Arg	Arg	Ala	Thr	Leu	Leu	Ser	Ala	Arg	Gln	Gly	Met	Met	Ser	Ala
				20					25					30
Arg	Gly	Asp	Phe	Leu	Asn	Tyr	Ala	Leu	Ser	Leu	Met	Arg	Ser	His
				35					40					45
Asn	Asp	Glu	His	Ser	Asp	Val	Leu	Pro	Val	Leu	Asp	Val	Cys	Ser
				50					55					60
Leu	Lys	His	Val	Ala	Tyr	Val	Phe	Gln	Ala	Leu	Ile	Tyr	Trp	Ile
				65					70					75
Lys	Ala	Met	Asn	Gln	Gln	Thr	Thr	Leu	Asp	Thr	Pro	Gln	Leu	Glu
				80					85					90
Arg	Lys	Arg	Thr	Arg	Glu	Leu	Leu	Glu	Leu	Gly	Ile	Asp	Asn	Glu
				95					100					105
Asp	Ser	Glu	His	Glu	Asn	Asp	Asp	Asp	Thr	Asn	Gln	Ser	Ala	Thr
				110					115					120
Leu	Asn	Asp	Lys	Asp	Asp	Asp	Ser	Leu	Pro	Ala	Glu	Thr	Gly	Gln
				125					130					135
Asn	His	Pro	Phe	Phe	Arg	Arg	Ser	Asp	Ser	Met	Thr	Phe	Leu	Gly
				140					145					150
Cys	Ile	Pro	Pro	Asn	Pro	Phe	Glu	Val	Pro	Leu	Ala	Glu	Ala	Ile
				155					160					165
Pro	Leu	Ala	Asp	Gln	Pro	His	Leu	Leu	Gln	Pro	Asn	Ala	Arg	Lys
				170					175					180
Glu	Asp	Leu	Phe	Gly	Arg	Pro	Ser	Gln	Gly	Leu	Tyr	Ser	Ser	Ser
				185					190					195
Ala	Ser	Ser	Gly	Lys	Cys	Leu	Met	Glu	Val	Thr	Val	Asp	Arg	Asn
				200					205					210
Cys	Leu	Glu	Val	Leu	Pro	Thr	Lys	Met	Ser	Tyr	Ala	Ala	Asn	Leu
				215					220					225
Lys	Asn	Val	Met	Asn	Met	Gln	Asn	Arg	Gln	Lys	Lys	Glu	Gly	Glu
				230					235					240
Glu	Gln	Pro	Val	Leu	Pro	Glu	Glu	Thr	Glu	Ser	Ser	Lys	Pro	Gly
				245					250					255
Pro	Ser	Ala	His	Asp	Leu	Ala	Ala	Gln	Leu	Lys	Ser	Ser	Leu	Leu
				260					265					270
Ala	Glu	Ile	Gly	Leu	Thr	Glu	Ser	Glu	Gly	Pro	Pro	Leu	Thr	Ser
				275					280					285
Phe	Arg	Pro	Gln	Cys	Ser	Phe	Met	Gly	Met	Val	Ile	Ser	His	Asp
				290					295					300
Met	Leu	Leu	Gly	Arg	Trp	Arg	Leu	Ser	Leu	Glu	Leu	Phe	Gly	Arg
				305					310					315
Val	Phe	Met	Glu	Asp	Val	Gly	Ala	Glu	Pro	Gly	Ser	Ile	Leu	Thr
				320					325					330
Glu	Leu	Gly	Gly	Phe	Glu	Val	Lys	Glu	Ser	Lys	Phe	Arg	Arg	Glu
				335					340					345
Met	Glu	Lys	Leu	Arg	Asn	Gln	Gln	Ser	Arg	Asp	Leu	Ser	Leu	Glu
				350					355					360
Val	Lys	Val	Asp	Arg	Asp	Arg	Asp	Leu	Leu	Ile	Gln	Gln	Thr	Met
				365					370					375
Arg	Gln	Leu	Asn	Asn	His	Phe	Gly	Arg	Arg	Cys	Ala	Thr	Thr	Pro
				380					385					390
Met	Ala	Val	His	Arg	Val	Lys	Val	Thr	Phe	Lys	Asp	Glu	Pro	Gly
				395					400					405

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Glu Gly Ser Gly Val Ala Arg Ser Phe Tyr Thr Ala Ile Ala Gln
 410 415 420
 Ala Phe Leu Ser Asn Glu Lys Leu Pro Asn Leu Glu Cys Ile Gln
 425 430 435
 Asn Ala Asn Lys Gly Thr His Thr Ser Leu Met Gln Arg Leu Arg
 440 445 450
 Asn Arg Gly Glu Arg Asp Arg Glu Arg Glu Arg Glu Arg Glu Met
 455 460 465
 Arg Arg Ser Ser Gly Leu Arg Ala Gly Ser Arg Arg Asp Arg Asp
 470 475 480
 Arg Asp Phe Arg Arg Gln Leu Ser Ile Asp Thr Arg Pro Phe Arg
 485 490 495
 Pro Ala Ser Glu Gly Asn Pro Ser Asp Asp Pro Glu Pro Leu Pro
 500 505 510
 Ala His Arg Gln Ala Leu Gly Glu Arg Leu Tyr Pro Arg Val Gln
 515 520 525
 Ala Met Gln Pro Ala Phe Ala Ser Lys Ile Thr Gly Met Leu Leu
 530 535 540
 Glu Leu Ser Pro Ala Gln Leu Leu Leu Leu Leu Ala Ser Glu Asp
 545 550 555
 Ser Leu Arg Ala Arg Val Asp Glu Ala Met Glu Leu Ile Ile Ala
 560 565 570
 His Gly Arg Glu Asn Gly Ala Asp Ser Ile Leu Asp Leu Gly Leu
 575 580 585
 Val Asp Ser Ser Glu Lys Val Gln Gln Glu Asn Arg Lys Arg His
 590 595 600
 Gly Ser Ser Arg Ser Val Val Asp Met Asp Leu Asp Asp Thr Asp
 605 610 615
 Asp Gly Asp Asp Asn Ala Pro Leu Phe Tyr Gln Pro Gly Lys Arg
 620 625 630
 Gly Phe Tyr Thr Pro Arg Pro Gly Lys Asn Thr Glu Ala Arg Leu
 635 640 645
 Asn Cys Phe Arg Asn Ile Gly Arg Ile Leu Gly Leu Cys Leu Leu
 650 655 660
 Gln Asn Glu Leu Cys Pro Ile Thr Leu Asn Arg His Val Ile Lys
 665 670 675
 Val Leu Leu Gly Arg Lys Val Asn Trp His Asp Phe Ala Phe Phe
 680 685 690
 Asp Pro Val Met Tyr Glu Ser Leu Arg Gln Leu Ile Leu Ala Ser
 695 700 705
 Gln Ser Ser Asp Ala Asp Ala Val Phe Ser Ala Met Asp Leu Ala
 710 715 720
 Phe Ala Ile Asp Leu Cys Lys Glu Glu Gly Gly Gln Val Glu
 725 730 735
 Leu Ile Pro Asn Gly Val Asn Ile Pro Val Thr Pro Gln Asn Val
 740 745 750
 Tyr Glu Tyr Val Arg Lys Tyr Ala Glu His Arg Met Leu Val Val
 755 760 765
 Ala Glu Gln Pro Leu His Ala Met Arg Lys Gly Leu Leu Asp Val
 770 775 780
 Leu Pro Lys Asn Ser Leu Glu Asp Leu Thr Ala Glu Asp Phe Arg
 785 790 795
 Leu Leu Val Asn Gly Cys Gly Glu Val Asn Val Gln Met Leu Ile
 800 805 810
 Ser Phe Thr Ser Phe Asp Asp Glu Ser Gly Glu Asn Ala Glu Lys

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815	820	825
Leu Leu Gln Phe Lys Arg Trp Phe Trp Ser Ile Val Glu Lys Met		
830	835	840
Ser Met Thr Glu Arg Gln Asp Leu Val Tyr Phe Trp Thr Ser Ser		
845	850	855
Pro Ser Leu Pro Ala Ser Glu Glu Gly Phe Gln Pro Met Pro Ser		
860	865	870
Ile Thr Ile Arg Pro Pro Asp Asp Gln His Leu Pro Thr Ala Asn		
875	880	885
Thr Cys Ile Ser Arg Leu Tyr Val Pro Leu Tyr Ser Ser Lys Gln		
890	895	900
Ile Leu Lys Gln Lys Leu Leu Ala Ile Lys Thr Lys Asn Phe		
905	910	915
Gly Phe Val		

<210> 6

<211> 324

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 2603810CD1

<400> 6

Met Gly Pro Trp Gly Glu Pro Glu Leu Leu Val Trp Arg Pro Glu			
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Ala Val Ala Ser Glu Pro Pro Val Pro Val Gly Leu Glu Val Lys			
20	25	30	
Leu Gly Ala Leu Val Leu Leu Leu Val Leu Thr Leu Leu Cys Ser			
35	40	45	
Leu Val Pro Ile Cys Val Leu Arg Arg Pro Gly Ala Asn His Glu			
50	55	60	
Gly Ser Ala Ser Arg Gln Lys Ala Leu Ser Leu Val Ser Cys Phe			
65	70	75	
Ala Gly Gly Val Phe Leu Ala Thr Cys Leu Leu Asp Leu Leu Pro			
80	85	90	
Asp Tyr Leu Ala Ala Ile Asp Glu Ala Leu Ala Ala Leu His Val			
95	100	105	
Thr Leu Gln Phe Pro Leu Gln Glu Phe Ile Leu Ala Met Gly Phe			
110	115	120	
Phe Leu Val Leu Val Met Glu Gln Ile Thr Leu Ala Tyr Lys Glu			
125	130	135	
Gln Ser Gly Pro Ser Pro Leu Glu Glu Thr Arg Ala Leu Leu Gly			
140	145	150	
Thr Val Asn Gly Gly Pro Gln His Trp His Asp Gly Pro Gly Val			
155	160	165	
Pro Gln Ala Ser Gly Ala Pro Ala Thr Pro Ser Ala Leu Arg Ala			
170	175	180	
Cys Val Leu Val Phe Ser Leu Ala Leu His Ser Val Phe Glu Gly			
185	190	195	
Leu Ala Val Gly Leu Gln Arg Asp Arg Ala Arg Ala Met Glu Leu			
200	205	210	
Cys Leu Ala Leu Leu His Lys Gly Ile Leu Ala Val Ser Leu			

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215	220	225
Ser Leu Arg Leu Leu Gln Ser His Leu Arg Ala Gln Val Val Ala		
230	235	240
Gly Cys Gly Ile Leu Phe Ser Cys Met Thr Pro Leu Gly Ile Gly		
245	250	255
Leu Gly Ala Ala Leu Ala Glu Ser Ala Gly Pro Leu His Gln Leu		
260	265	270
Ala Gln Ser Val Leu Glu Gly Met Ala Ala Gly Thr Phe Leu Tyr		
275	280	285
Ile Thr Phe Leu Glu Ile Leu Pro Gln Glu Leu Ala Ser Ser Glu		
290	295	300
Gln Arg Ile Leu Lys Val Ile Leu Leu Leu Ala Gly Phe Ala Leu		
305	310	315
Leu Thr Gly Leu Leu Phe Ile Gln Ile		
320		

<210> 7
<211> 185
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 2715761CD1

<400> 7

Met Thr Thr Pro Asn Lys Thr Pro Pro Gly Ala Asp Pro Lys Gln			
1	5	10	15
Leu Glu Arg Thr Gly Thr Val Arg Glu Ile Gly Ser Gln Ala Val			
20	25	30	
Trp Ser Leu Ser Ser Cys Lys Pro Gly Phe Gly Val Asp Gln Leu			
35	40	45	
Arg Asp Asp Asn Leu Glu Thr Tyr Trp Gln Ser Asp Gly Ser Gln			
50	55	60	
Pro His Leu Val Asn Ile Gln Phe Arg Arg Lys Thr Thr Val Lys			
65	70	75	
Thr Leu Cys Ile Tyr Ala Asp Tyr Lys Ser Asp Glu Ser Tyr Thr			
80	85	90	
Pro Ser Lys Ile Ser Val Arg Val Gly Asn Asn Phe His Asn Leu			
95	100	105	
Gln Glu Ile Arg Gln Leu Glu Leu Val Glu Pro Ser Gly Trp Ile			
110	115	120	
His Val Pro Leu Thr Asp Asn His Lys Lys Pro Thr Arg Thr Phe			
125	130	135	
Met Ile Gln Ile Ala Val Leu Ala Asn His Gln Asn Gly Arg Asp			
140	145	150	
Thr His Met Arg Gln Ile Lys Ile Tyr Thr Pro Val Glu Glu Ser			
155	160	165	
Ser Ile Gly Lys Phe Pro Arg Cys Thr Thr Ile Asp Phe Met Met			
170	175	180	
Tyr Arg Ser Ile Arg			
185			

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<210> 8
 <211> 445
 <212> PRT
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> Incyte ID No: 3255641CD1

<400> 8

Met	Leu	Ala	Ser	Tyr	Gly	Leu	Ala	Tyr	Ser	Leu	Met	Lys	Phe	Phe
1						5				10				15
Thr	Gly	Pro	Met	Ser	Asp	Phe	Lys	Asn	Val	Gly	Leu	Val	Phe	Val
			20						25					30
Asn	Ser	Lys	Arg	Asp	Arg	Thr	Lys	Ala	Val	Leu	Cys	Met	Val	Val
			35						40					45
Ala	Gly	Ala	Ile	Ala	Ala	Val	Phe	His	Thr	Leu	Ile	Ala	Tyr	Ser
			50						55					60
Asp	Leu	Gly	Tyr	Tyr	Ile	Ile	Asn	Lys	Leu	His	His	Val	Asp	Glu
			65						70					75
Ser	Val	Gly	Ser	Lys	Thr	Arg	Arg	Ala	Phe	Leu	Tyr	Leu	Ala	Ala
			80						85					90
Phe	Pro	Phe	Met	Asp	Ala	Met	Ala	Trp	Thr	His	Ala	Gly	Ile	Leu
			95						100					105
Leu	Lys	His	Lys	Tyr	Ser	Phe	Leu	Val	Gly	Cys	Ala	Ser	Ile	Ser
			110						115					120
Asp	Val	Ile	Ala	Gln	Val	Val	Phe	Val	Ala	Ile	Leu	Leu	His	Ser
			125						130					135
His	Leu	Glu	Cys	Arg	Glu	Pro	Leu	Leu	Ile	Pro	Ile	Leu	Ser	Leu
			140						145					150
Tyr	Met	Gly	Ala	Leu	Val	Arg	Cys	Thr	Thr	Leu	Cys	Leu	Gly	Tyr
			155						160					165
Tyr	Lys	Asn	Ile	His	Asp	Ile	Ile	Pro	Asp	Arg	Ser	Gly	Pro	Glu
			170						175					180
Leu	Gly	Gly	Asp	Ala	Thr	Ile	Arg	Lys	Met	Leu	Ser	Phe	Trp	Trp
			185						190					195
Pro	Leu	Ala	Leu	Ile	Leu	Ala	Thr	Gln	Arg	Ile	Ser	Arg	Pro	Ile
			200						205					210
Val	Asn	Leu	Phe	Val	Ser	Arg	Asp	Leu	Gly	Gly	Ser	Ser	Ala	Ala
			215						220					225
Thr	Glu	Ala	Val	Ala	Ile	Leu	Thr	Ala	Thr	Tyr	Pro	Val	Gly	His
			230						235					240
Met	Pro	Tyr	Gly	Trp	Leu	Thr	Glu	Ile	Arg	Ala	Val	Tyr	Pro	Ala
			245						250					255
Phe	Asp	Lys	Asn	Asn	Pro	Ser	Asn	Lys	Leu	Val	Ser	Thr	Ser	Asn
			260						265					270
Thr	Val	Thr	Ala	Ala	His	Ile	Lys	Lys	Phe	Thr	Phe	Val	Cys	Met
			275						280					285
Ala	Leu	Ser	Leu	Thr	Leu	Cys	Phe	Val	Met	Phe	Trp	Thr	Pro	Asn
			290						295					300
Val	Ser	Glu	Lys	Ile	Leu	Ile	Asp	Ile	Ile	Gly	Val	Asp	Phe	Ala
			305						310					315
Phe	Ala	Glu	Leu	Cys	Val	Val	Pro	Leu	Arg	Ile	Phe	Ser	Phe	Phe
			320						325					330
Pro	Val	Pro	Val	Thr	Val	Arg	Ala	His	Leu	Thr	Gly	Trp	Leu	Met

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335	340	345
Thr Leu Lys Lys	Thr Phe Val Leu Ala Pro Ser Ser Val Leu Arg	
350	355	360
Ile Ile Val Leu Ile Ala Ser Leu Val Val	Leu Pro Tyr Leu Gly	
365	370	375
Val His Gly Ala	Thr Leu Gly Val Gly Ser Leu Leu Ala Gly Phe	
380	385	390
Val Gly Glu Ser	Thr Met Val Ala Ile Ala Ala Cys Tyr Val Tyr	
395	400	405
Arg Lys Gln Lys	Lys Lys Met Glu Asn Glu Ser Ala Thr Glu Gly	
410	415	420
Glu Asp Ser Ala Met	Thr Asp Met Pro Pro Thr Glu Glu Val Thr	
425	430	435
Asp Ile Val Glu Met Arg Glu Glu Asn Glu		
440	445	

<210> 9
<211> 73
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 3620391CD1

<400> 9		
Met Pro Arg Glu Arg Arg Glu Arg Asp Ala Lys Glu Arg Asp Thr		
1 5 10 15		
Met Lys Glu Asp Gly Gly Ala Glu Phe Ser Ala Arg Ser Arg Lys		
20 25 30		
Arg Lys Ala Asn Val Thr Val Phe Cys Arg Ile Gln Met Lys Lys		
35 40 45		
Trp Pro Lys Ser Thr Gly Arg Arg Trp Thr Ser Val Gly Ala Arg		
50 55 60		
Leu Gly Arg Met Met Gln Ser Val Gln Ala Pro Ala Pro		
65 70		

<210> 10
<211> 288
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 3969860CD1

<400> 10		
Met Ala Ala Leu Phe Gln Glu Ala Ser Ser Cys Pro Val Cys Ser		
1 5 10 15		
Asp Tyr Leu Glu Lys Pro Met Ser Leu Glu Cys Gly Cys Ala Val		
20 25 30		
Cys Leu Lys Cys Ile Asn Ser Leu Gln Lys Glu Pro His Gly Glu		
35 40 45		

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Asp	Leu	Leu	Cys	Cys	Cys	Ser	Ser	Met	Val	Ser	Arg	Lys	Asn	Lys
								50		55				60
Ile	Arg	Arg	Asn	Arg	Gln	Leu	Glu	Arg	Leu	Ala	Ser	His	Ile	Lys
								65		70				75
Glu	Leu	Glu	Pro	Lys	Leu	Lys	Lys	Ile	Leu	Gln	Met	Asn	Pro	Arg
								80		85				90
Met	Arg	Lys	Phe	Gln	Val	Asp	Met	Thr	Leu	Asp	Ala	Asn	Thr	Ala
								95		100				105
Asn	Asn	Phe	Leu	Leu	Ile	Ser	Asp	Asp	Leu	Arg	Ser	Val	Arg	Ser
								110		115				120
Gly	Arg	Ile	Arg	Gln	Asn	Arg	Gln	Asp	Leu	Ala	Glu	Arg	Phe	Asp
								125		130				135
Val	Ser	Val	Cys	Ile	Leu	Gly	Ser	Pro	Arg	Phe	Thr	Cys	Gly	Arg
								140		145				150
His	Cys	Trp	Glu	Val	Asp	Val	Gly	Thr	Ser	Thr	Glu	Trp	Asp	Leu
								155		160				165
Gly	Val	Cys	Arg	Glu	Ser	Val	His	Arg	Lys	Gly	Arg	Ile	Gln	Leu
								170		175				180
Thr	Thr	Glu	Leu	Gly	Phe	Trp	Thr	Val	Ser	Leu	Arg	Asp	Gly	Gly
								185		190				195
Arg	Leu	Ser	Ala	Ser	Thr	Val	Pro	Leu	Thr	Phe	Leu	Phe	Val	Asp
								200		205				210
Arg	Lys	Leu	Gln	Arg	Val	Gly	Ile	Phe	Leu	Asp	Met	Gly	Met	Gln
								215		220				225
Asn	Val	Ser	Phe	Phe	Asp	Ala	Glu	Ser	Gly	Ser	His	Val	Tyr	Thr
								230		235				240
Phe	Arg	Ser	Val	Ser	Ala	Glu	Glu	Pro	Leu	Arg	Pro	Phe	Leu	Ala
								245		250				255
Pro	Ser	Val	Pro	Pro	Asn	Gly	Asp	Gln	Gly	Val	Leu	Ser	Ile	Cys
								260		265				270
Pro	Leu	Met	Asn	Ser	Gly	Thr	Thr	Asp	Ala	Pro	Val	Arg	Pro	Gly
								275		280				285
Glu	Ala	Lys												

<210> 11
<211> 98
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 4286006CD1

<400> 11
Met Ala Lys Phe Gly Val His Arg Ile Leu Leu Ala Ile Ser
1 5 10 15
Leu Thr Lys Cys Leu Glu Ser Thr Lys Leu Leu Ala Asp Leu Lys
20 25 30
Lys Cys Gly Asp Leu Glu Cys Glu Ala Leu Ile Asn Arg Val Ser
35 40 45
Ala Met Arg Asp Tyr Arg Gly Pro Asp Cys Arg Tyr Leu Asn Phe
50 55 60
Thr Lys Gly Glu Glu Ile Ser Val Tyr Val Lys Leu Ala Gly Asp

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65	70	75
Arg Glu Asp Leu Trp Ala Gly Ser Lys Gly	Lys Glu Phe Gly Tyr	
80	85	90
Phe Pro Arg Asp Ala Val Gln Ile		
95		

<210> 12

<211> 549

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 4325626CD1

<400> 12

Met Asp Val Val Glu Val Ala Gly Ser Trp Trp Ala Gln Glu Arg			
1	5	10	15
Glu Asp Ile Ile Met Lys Tyr Glu Lys Gly His Arg Ala Gly Leu			
20		25	30
Pro Glu Asp Lys Gly Pro Lys Pro Phe Arg Ser Tyr Asn Asn Asn			
35		40	45
Val Asp His Leu Gly Ile Val His Glu Thr Glu Leu Pro Pro Leu			
50		55	60
Thr Ala Arg Glu Ala Lys Gln Ile Arg Arg Glu Ile Ser Arg Lys			
65		70	75
Ser Lys Trp Val Asp Met Leu Gly Asp Trp Glu Lys Tyr Lys Ser			
80		85	90
Ser Arg Lys Leu Ile Asp Arg Ala Tyr Lys Gly Met Pro Met Asn			
95		100	105
Ile Arg Gly Pro Met Trp Ser Val Leu Leu Asn Thr Glu Glu Met			
110		115	120
Lys Leu Lys Asn Pro Gly Arg Tyr Gln Ile Met Lys Glu Lys Gly			
125		130	135
Lys Arg Ser Ser Glu His Ile Gln Arg Ile Asp Arg Asp Val Ser			
140		145	150
Gly Thr Leu Arg Lys His Ile Phe Phe Arg Asp Arg Tyr Gly Thr			
155		160	165
Lys Gln Arg Glu Leu Leu His Ile Leu Leu Ala Tyr Glu Glu Tyr			
170		175	180
Asn Pro Glu Val Gly Tyr Cys Arg Asp Leu Ser His Ile Ala Ala			
185		190	195
Leu Phe Leu Leu Tyr Leu Pro Glu Glu Asp Ala Phe Trp Ala Leu			
200		205	210
Val Gln Leu Leu Ala Ser Glu Arg His Ser Leu Gln Gly Phe His			
215		220	225
Ser Pro Asn Gly Gly Thr Val Gln Gly Leu Gln Asp Gln Gln Glu			
230		235	240
His Val Val Ala Thr Ser Gln Pro Lys Thr Met Gly His Gln Asp			
245		250	255
Lys Lys Asp Leu Cys Gly Gln Cys Ser Pro Leu Gly Cys Leu Ile			
260		265	270
Arg Ile Leu Ile Asp Gly Ile Ser Leu Gly Leu Thr Leu Arg Leu			
275		280	285
Trp Asp Val Tyr Leu Val Glu Gly Glu Gln Ala Leu Met Pro Ile			

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290	295	300
Thr Arg Ile Ala Phe Lys Val Gln Gln	Lys Arg Leu Thr Lys	Thr
305	310	315
Ser Arg Cys Gly Pro Trp Ala Arg Phe	Cys Asn Arg Phe Val	Asp
320	325	330
Thr Trp Ala Arg Asp Glu Asp Thr Val	Leu Lys His Leu Arg	Ala
335	340	345
Ser Met Lys Lys Leu Thr Arg Lys Gln	Gly Asp Leu Pro Pro	Pro
350	355	360
Ala Lys Pro Glu Gln Gly Ser Ser Ala	Ser Arg Pro Val Pro	Ala
365	370	375
Ser Arg Gly Gly Lys Thr Leu Cys Lys	Gly Asp Arg Gln Ala	Pro
380	385	390
Pro Gly Pro Pro Ala Arg Phe Pro Arg	Pro Ile Trp Ser Ala	Ser
395	400	405
Pro Pro Arg Ala Pro Arg Ser Ser Thr	Pro Cys Pro Gly Gly	Ala
410	415	420
Val Arg Glu Asp Thr Tyr Pro Val Gly	Thr Gln Gly Val Pro	Ser
425	430	435
Pro Ala Leu Ala Gln Gly Gly Pro Gln	Gly Ser Trp Arg Phe	Leu
440	445	450
Gln Trp Asn Ser Met Pro Arg Leu Pro	Thr Asp Leu Asp Val	Glu
455	460	465
Gly Pro Trp Phe Arg His Tyr Asp Phe	Arg Gln Ser Cys Trp	Val
470	475	480
Arg Ala Ile Ser Gln Glu Asp Gln Leu	Ala Pro Cys Trp Gln	Ala
485	490	495
Glu His Pro Ala Glu Arg Val Arg Ser	Ala Phe Ala Ala Pro	Ser
500	505	510
Thr Asp Ser Asp Gln Gly Thr Pro Phe	Arg Ala Arg Asp Glu	Gln
515	520	525
Pro Cys Ala Pro Thr Ser Gly Pro Cys	Leu Cys Gly Leu His	Leu
530	535	540
Glu Ser Ser Gln Phe Pro Pro Gly Phe		
545		

<210> 13
 <211> 95
 <212> PRT
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> Incyte ID No: 1438978CD1

<400> 13

Met Ser Phe Leu Leu Pro Lys Leu Thr Ser	Lys Lys Glu Val Asp		
1	5	10	15
Gln Ala Ile Lys Ser Thr Ala Glu Lys Val	Leu Val Leu Arg Phe		
20	25	30	
Gly Arg Asp Glu Asp Pro Val Cys Leu Gln	Leu Asp Asp Ile Leu		
35	40	45	
Ser Lys Thr Ser Ser Asp Leu Ser Lys Met	Ala Ala Ile Tyr Leu		
50	55	60	

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Val Asp Val Asp Gln Thr Ala Val Tyr Thr Gln Tyr Phe Asp Ile
 65 70 75
 Ser Tyr Ile Pro Ser Thr Val Phe Phe Phe Asn Gly Gln His Met
 80 85 90
 Lys Val Asp Tyr Gly
 95

<210> 14
 <211> 445
 <212> PRT
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> Incyte ID No: 2024773CD1

<400> 14

Met Ala Ala Pro Glu Glu Arg Asp Leu Thr Gln Glu Gln Thr Glu	
1 5 10 15	
Lys Leu Leu Gln Phe Gln Asp Leu Thr Gly Ile Glu Ser Met Asp	
20 25 30	
Gln Cys Arg His Thr Leu Glu Gln His Asn Trp Asn Ile Glu Ala	
35 40 45	
Ala Val Gln Asp Arg Leu Asn Glu Gln Glu Gly Val Pro Ser Val	
50 55 60	
Phe Asn Pro Pro Ser Arg Pro Leu Gln Val Asn Thr Ala Asp	
65 70 75	
His Arg Ile Tyr Ser Tyr Val Val Ser Arg Pro Gln Pro Arg Gly	
80 85 90	
Leu Leu Gly Trp Gly Tyr Tyr Leu Ile Met Leu Pro Phe Arg Phe	
95 100 105	
Thr Tyr Tyr Thr Ile Leu Asp Ile Phe Arg Phe Ala Leu Arg Phe	
110 115 120	
Ile Arg Pro Asp Pro Arg Ser Arg Val Thr Asp Pro Val Gly Asp	
125 130 135	
Ile Val Ser Phe Met His Ser Phe Glu Glu Lys Tyr Gly Arg Ala	
140 145 150	
His Pro Val Phe Tyr Gln Gly Thr Tyr Ser Gln Ala Leu Asn Asp	
155 160 165	
Ala Lys Arg Glu Leu Arg Phe Leu Leu Val Tyr Leu His Gly Asp	
170 175 180	
Asp His Gln Asp Ser Asp Glu Phe Cys Arg Asn Thr Leu Cys Ala	
185 190 195	
Pro Glu Val Ile Ser Leu Ile Asn Thr Arg Met Leu Phe Trp Ala	
200 205 210	
Cys Ser Thr Asn Lys Pro Glu Gly Tyr Arg Val Ser Gln Ala Leu	
215 220 225	
Arg Glu Asn Thr Tyr Pro Phe Leu Ala Met Ile Met Leu Lys Asp	
230 235 240	
Arg Arg Met Thr Val Val Gly Arg Leu Glu Gly Leu Ile Gln Pro	
245 250 255	
Asp Asp Leu Ile Asn Gln Leu Thr Phe Ile Met Asp Ala Asn Gln	
260 265 270	
Thr Tyr Leu Val Ser Glu Arg Leu Glu Arg Glu Glu Arg Asn Gln	
275 280 285	

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Thr	Gln	Val	Leu	Arg	Gln	Gln	Gln	Asp	Glu	Ala	Tyr	Leu	Ala	Ser
					290				295					300
Leu	Arg	Ala	Asp	Gln	Glu	Lys	Glu	Arg	Lys	Lys	Arg	Glu	Glu	Arg
					305				310					315
Glu	Arg	Lys	Arg	Arg	Lys	Glu	Glu	Glu	Val	Gln	Gln	Gln	Lys	Leu
					320				325					330
Ala	Glu	Glu	Arg	Arg	Arg	Gln	Asn	Leu	Gln	Glu	Glu	Lys	Glu	Arg
					335				340					345
Lys	Leu	Glu	Cys	Leu	Pro	Pro	Glu	Pro	Ser	Pro	Asp	Asp	Pro	Glu
					350				355					360
Ser	Val	Lys	Ile	Ile	Phe	Lys	Leu	Pro	Asn	Asp	Ser	Arg	Val	Glu
					365				370					375
Arg	Arg	Phe	His	Phe	Ser	Gln	Ser	Leu	Thr	Val	Ile	His	Asp	Phe
					380				385					390
Leu	Phe	Ser	Leu	Lys	Glu	Ser	Pro	Glu	Lys	Phe	Gln	Ile	Glu	Ala
					395				400					405
Asn	Phe	Pro	Arg	Arg	Val	Leu	Pro	Cys	Ile	Pro	Ser	Glu	Glu	Trp
					410				415					420
Pro	Asn	Pro	Pro	Thr	Leu	Gln	Glu	Ala	Gly	Leu	Ser	His	Thr	Glu
					425				430					435
Val	Leu	Phe	Val	Gln	Asp	Leu	Thr	Asp	Glu					
					440				445					

<210> 15
<211> 219
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 3869790CD1

Met	Glu	Tyr	Leu	Ser	Ala	Leu	Asn	Pro	Ser	Asp	Leu	Leu	Arg	Ser
1					5				10					15
Val	Ser	Asn	Ile	Ser	Ser	Glu	Phe	Gly	Arg	Arg	Val	Trp	Thr	Ser
						20			25					30
Ala	Pro	Pro	Pro	Gln	Arg	Pro	Phe	Arg	Val	Cys	Asp	His	Lys	Arg
					35				40					45
Thr	Ile	Arg	Lys	Gly	Leu	Thr	Ala	Ala	Thr	Arg	Gln	Glu	Leu	Leu
					50				55					60
Ala	Lys	Ala	Leu	Glu	Thr	Leu	Leu	Leu	Asn	Gly	Val	Leu	Thr	Leu
					65				70					75
Val	Leu	Glu	Glu	Asp	Gly	Thr	Ala	Val	Asp	Ser	Glu	Asp	Phe	Phe
					80				85					90
Gln	Leu	Leu	Glu	Asp	Asp	Thr	Cys	Leu	Met	Val	Leu	Gln	Ser	Gly
						95				100				105
Gln	Ser	Trp	Ser	Pro	Thr	Arg	Ser	Gly	Val	Leu	Ser	Tyr	Gly	Leu
					110				115					120
Gly	Arg	Glu	Arg	Pro	Lys	His	Ser	Lys	Asp	Ile	Ala	Arg	Phe	Thr
					125				130					135
Phe	Asp	Val	Tyr	Lys	Gln	Asn	Pro	Arg	Asp	Leu	Phe	Gly	Ser	Leu
					140				145					150
Asn	Val	Lys	Ala	Thr	Phe	Tyr	Gly	Leu	Tyr	Ser	Met	Ser	Cys	Asp

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155	160	165
Phe Gln Gly Leu Gly Pro Lys Lys Val	Leu Arg Glu Leu Leu Arg	
170	175	180
Trp Thr Ser Thr Leu Leu Gln Gly Leu	Gly His Met Leu Leu Gly	
185	190	195
Ile Ser Ser Thr Leu Arg His Ala Val	Glu Gly Ala Glu Gln Trp	
200	205	210
Gln Gln Lys Gly Arg Leu His Ser Tyr		
215		

<210> 16

<211> 439

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 001273CD1

<400> 16

Met Ala Ala Ala Arg Cys Trp Arg Pro Leu Leu Arg Gly Pro Arg			
1	5	10	15
Leu Ser Leu His Thr Ala Ala Asn Ala Ala Thr Ala Thr Glu			
20	25	30	
Thr Thr Cys Gln Asp Val Ala Ala Thr Pro Val Ala Arg Tyr Pro			
35	40	45	
Pro Ile Val Ala Ser Met Thr Ala Asp Ser Lys Ala Ala Arg Leu			
50	55	60	
Arg Arg Ile Glu Arg Trp Gln Ala Thr Val His Ala Ala Glu Ser			
65	70	75	
Val Asp Glu Lys Leu Arg Ile Leu Thr Lys Met Gln Phe Met Lys			
80	85	90	
Tyr Met Val Tyr Pro Gln Thr Phe Ala Leu Asn Ala Asp Arg Trp			
95	100	105	
Tyr Gln Tyr Phe Thr Lys Thr Val Phe Leu Ser Gly Leu Pro Pro			
110	115	120	
Arg Pro Ser Glu Pro Glu Pro Glu Pro Glu Pro Glu Pro			
125	130	135	
Ala Leu Asp Leu Ala Ala Leu Arg Ala Val Ala Cys Asp Cys Leu			
140	145	150	
Leu Gln Glu His Phe Tyr Leu Arg Arg Arg Arg Arg Val His Arg			
155	160	165	
Tyr Glu Glu Ser Glu Val Ile Ser Leu Pro Phe Leu Asp Gln Leu			
170	175	180	
Val Ser Thr Leu Val Gly Leu Leu Ser Pro His Asn Pro Ala Leu			
185	190	195	
Ala Ala Ala Ala Leu Asp Tyr Arg Cys Pro Val His Phe Tyr Trp			
200	205	210	
Val Arg Gly Glu Glu Ile Ile Pro Arg Gly His Arg Arg Gly Arg			
215	220	225	
Ile Asp Asp Leu Arg Tyr Gln Ile Asp Asp Lys Pro Asn Asn Gln			
230	235	240	
Ile Arg Ile Ser Lys Gln Leu Ala Glu Phe Val Pro Leu Asp Tyr			
245	250	255	
Ser Val Pro Ile Glu Ile Pro Thr Ile Lys Cys Lys Pro Asp Lys			

260	265	270
Leu Pro Leu Phe Lys Arg Gln Tyr Glu	Asn His Ile Phe Val	Gly
275	280	285
Ser Lys Thr Ala Asp Pro Cys Cys Tyr	Gly His Thr Gln Phe His	
290	295	300
Leu Leu Pro Asp Lys Leu Arg Arg Glu	Arg Leu Leu Arg Gln Asn	
305	310	315
Cys Ala Asp Gln Ile Glu Val Val Phe	Arg Ala Asn Ala Ile Ala	
320	325	330
Ser Leu Phe Ala Trp Thr Gly Ala Gln	Ala Met Tyr Gln Gly Phe	
335	340	345
Trp Ser Glu Ala Asp Val Thr Arg Pro	Phe Val Ser Gln Ala Val	
350	355	360
Ile Thr Asp Gly Lys Tyr Phe Ser Phe	Phe Cys Tyr Gln Leu Asn	
365	370	375
Thr Leu Ala Leu Thr Thr Gln Ala Asp	Gln Asn Asn Pro Arg Lys	
380	385	390
Asn Ile Cys Trp Gly Thr Gln Ser Lys	Pro Leu Tyr Glu Thr Ile	
395	400	405
Glu Asp Asn Asp Val Lys Gly Phe Asn	Asp Asp Val Leu Leu Gln	
410	415	420
Ile Val His Phe Leu Leu Asn Arg Pro	Lys Glu Glu Lys Ser Gln	
425	430	435
Leu Leu Glu Asn		

<210> 17
<211> 526
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 411831CD1

<400> 17		
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Ser Ser Ala Ala Pro Ser Ala Gly Gly	Ser Ser Ser Gly Thr Thr	
20 25 30		
Thr Thr Thr Thr Thr Gly Gly Ile Leu Ile Gly Asp Arg		
35 40 45		
Leu Tyr Ser Glu Val Ser Leu Thr Ile Asp His Ser Leu Ile Pro		
50 55 60		
Glu Glu Arg Leu Ser Pro Thr Pro Ser Met Gln Asp Gly Leu Asp		
65 70 75		
Leu Pro Ser Glu Thr Asp Leu Arg Ile Leu Gly Cys Glu Leu Ile		
80 85 90		
Gln Ala Ala Gly Ile Leu Leu Arg Leu Pro Gln Val Ala Met Ala		
95 100 105		
Thr Gly Gln Val Leu Phe His Arg Phe Phe Tyr Ser Lys Ser Phe		
110 115 120		
Val Lys His Ser Phe Glu Ile Val Ala Met Ala Cys Ile Asn Leu		
125 130 135		
Ala Ser Lys Ile Glu Glu Ala Pro Arg Arg Ile Arg Asp Val Ile		

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140	145	150
Asn Val Phe His His Leu Arg Gln Leu Arg	Gly Lys Arg Thr	Pro
155	160	165
Ser Pro Leu Ile Leu Asp Gln Asn Tyr	Ile Asn Thr Lys Asn	Gln
170	175	180
Val Ile Lys Ala Glu Arg Arg Val Leu	Lys Glu Leu Gly Phe	Cys
185	190	195
Val His Val Lys His Pro His Lys Ile	Ile Val Met Tyr Leu	Gln
200	205	210
Val Leu Glu Cys Glu Arg Asn Gln Thr	Leu Val Gln Thr Ala	Trp
215	220	225
Asn Tyr Met Asn Asp Ser Leu Arg Thr	Asn Val Phe Val Arg	Phe
230	235	240
Gln Pro Glu Thr Ile Ala Cys Ala Cys	Ile Tyr Leu Ala Ala	Arg
245	250	255
Ala Leu Gln Ile Pro Leu Pro Thr Arg	Pro His Trp Phe Leu	Leu
260	265	270
Phe Gly Thr Thr Glu Glu Glu Ile Gln	Glu Ile Cys Ile Glu	Thr
275	280	285
Leu Arg Leu Tyr Thr Arg Lys Lys Pro	Asn Tyr Glu Leu Leu	Glu
290	295	300
Lys Glu Val Glu Lys Arg Lys Val Ala	Leu Gln Glu Ala Lys	Leu
305	310	315
Lys Ala Lys Gly Leu Asn Pro Asp Gly	Thr Pro Ala Leu Ser	Thr
320	325	330
Leu Gly Gly Phe Ser Pro Ala Ser Lys	Pro Ser Ser Pro Arg	Glu
335	340	345
Val Lys Ala Glu Glu Lys Ser Pro Ile	Ser Ile Asn Val Lys	Thr
350	355	360
Val Lys Lys Glu Pro Glu Asp Arg Gln	Gln Ala Ser Lys Ser	Pro
365	370	375
Tyr Asn Gly Val Arg Lys Asp Ser Lys	Arg Ser Arg Asn Ser	Arg
380	385	390
Ser Ala Ser Arg Ser Arg Ser Arg Thr	Arg Ser Arg Ser Arg	Ser
395	400	405
His Thr Pro Arg Arg His Tyr Asn Asn	Arg Arg Ser Arg Ser	Gly
410	415	420
Thr Tyr Ser Ser Arg Ser Arg Ser Arg	Ser Arg Ser His Ser	Glu
425	430	435
Ser Pro Arg Arg His His Asn His Gly	Ser Pro His Leu Lys	Ala
440	445	450
Lys His Thr Arg Asp Asp Leu Lys Ser	Ser Asn Arg His Gly	His
455	460	465
Lys Arg Lys Lys Ser Arg Ser Arg Ser	Gln Ser Lys Ser Arg	Asp
470	475	480
His Ser Asp Ala Ala Lys Lys His Arg	His Glu Arg Gly His	His
485	490	495
Arg Asp Arg Arg Glu Arg Ser Arg Ser	Phe Glu Arg Ser His	Lys
500	505	510
Ser Lys His His Gly Gly Ser Arg Ser	Gly His Gly Arg His	Arg
515	520	525

Arg

<210> 18

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<211> 298
 <212> PRT
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> Incyte ID No: 1520835CD1

<400> 18

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Gly	Pro	Arg	Ser	Leu	Gly	Ser	Pro	Val	Leu	Gly	Leu	Asp	Thr	Cys	
										40					45
Arg	Ala	Trp	Asp	His	Val	Asp	Gly	Gln	Ile	Leu	Gly	Gln	Leu	Arg	
										55					60
Pro	Leu	Thr	Glu	Glu	Glu	Glu	Glu	Gly	Ala	Gly	Ala	Thr	Leu		
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Ser	Arg	Gly	Pro	Ala	Phe	Pro	Gly	Met	Gly	Ser	Glu	Glu	Leu	Arg	
										85					90
Leu	Ala	Ser	Phe	Tyr	Asp	Trp	Pro	Leu	Thr	Ala	Glu	Val	Pro	Pro	
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Glu	Leu	Leu	Ala	Ala	Gly	Phe	Phe	His	Thr	Gly	His	Gln	Asp		
										110					120
Lys	Val	Arg	Cys	Phe	Phe	Cys	Tyr	Gly	Gly	Leu	Gln	Ser	Trp	Lys	
										125					135
Arg	Gly	Asp	Asp	Pro	Trp	Thr	Glu	His	Ala	Lys	Trp	Phe	Pro	Ser	
										140					150
Cys	Gln	Phe	Leu	Leu	Arg	Ser	Lys	Gly	Arg	Asp	Phe	Val	His	Ser	
										155					165
Val	Gln	Glu	Thr	His	Ser	Gln	Leu	Leu	Gly	Ser	Trp	Asp	Pro	Trp	
										170					180
Glu	Glu	Pro	Glu	Asp	Ala	Ala	Pro	Val	Ala	Pro	Ser	Val	Pro	Ala	
										185					195
Ser	Gly	Tyr	Pro	Glu	Leu	Pro	Thr	Pro	Arg	Arg	Glu	Val	Gln	Ser	
										200					210
Glu	Ser	Ala	Gln	Glu	Pro	Gly	Gly	Val	Ser	Pro	Ala	Glu	Ala	Gln	
										215					225
Arg	Ala	Trp	Trp	Val	Leu	Glu	Pro	Pro	Gly	Ala	Arg	Asp	Val	Glu	
										230					240
Ala	Gln	Leu	Arg	Arg	Leu	Gln	Glu	Glu	Arg	Thr	Cys	Lys	Val	Cys	
										245					255
Leu	Asp	Arg	Ala	Val	Ser	Ile	Val	Phe	Val	Pro	Cys	Gly	His	Leu	
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Val	Cys	Ala	Glu	Cys	Ala	Pro	Gly	Leu	Gln	Leu	Cys	Pro	Ile	Cys	
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Arg	Ala	Pro	Val	Arg	Ser	Arg	Val	Arg	Thr	Phe	Leu	Ser			
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<210> 19
 <211> 249
 <212> PRT
 <213> Homo sapiens

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<220>
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 <223> Incyte ID No: 1902803CD1

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 35 40 45
 Ala Thr Glu Tyr Met Ser Ser Ala Arg Ser Leu Ser Ser Glu Glu
 50 55 60
 Lys Leu Ala Leu Leu Lys Gln Ile Gln Glu Ala Tyr Gly Lys Cys
 65 70 75
 Lys Glu Phe Gly Asp Asp Lys Val Gln Leu Ala Met Gln Thr Tyr
 80 85 90
 Glu Met Val Asp Lys His Ile Arg Arg Leu Asp Thr Asp Leu Ala
 95 100 105
 Arg Phe Glu Ala Asp Leu Lys Glu Lys Gln Ile Glu Ser Ser Asp
 110 115 120
 Tyr Asp Ser Ser Ser Lys Gly Lys Lys Gly Arg Thr Gln
 125 130 135
 Lys Glu Lys Lys Ala Ala Arg Ala Arg Ser Lys Gly Lys Asn Ser
 140 145 150
 Asp Glu Glu Ala Pro Lys Thr Ala Gln Lys Lys Leu Lys Leu Val
 155 160 165
 Arg Thr Ser Pro Glu Tyr Gly Met Pro Ser Val Thr Phe Gly Ser
 170 175 180
 Val His Pro Ser Asp Val Leu Asp Met Pro Val Asp Pro Asn Glu
 185 190 195
 Pro Thr Tyr Cys Leu Cys His Gln Val Ser Tyr Gly Glu Met Ile
 200 205 210
 Gly Cys Asp Asn Pro Asp Cys Ser Ile Glu Trp Phe His Phe Ala
 215 220 225
 Cys Val Gly Leu Thr Thr Lys Pro Arg Gly Lys Trp Phe Cys Pro
 230 235 240
 Arg Cys Ser Gln Glu Arg Lys Lys Lys
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<210> 20
 <211> 1748
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> Incyte ID No: 1342011CB1

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 gccgcatcg ttgctgctcc ccagagacag acctgggccc ttccctctgg gactccaaat 180
 ctggacgggg ttcctggctt gctgtgggc atgttgaggc cggaggctgg gttgtgggg 240

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 ataagtccctt ggactgttcg cttccgggtt ctgagccctg gcgtcaggga gaaagggcat 420
 gtccagaaca atggccagaa ccaggcccg ccagctcggt cgggtacgg gggcggtgg 480
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 cctgttcggt ttagacccccc aaactggagg ggcatggag aaccgttagag cgcaggaacg 1680
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 aaaaaagg 1748

<210> 21
 <211> 1016
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> Incyte ID No: 1880041CB1

<400> 21

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 gagcatgaat acatccacaa cttcaagggtg ctgcaagcag ctttcaagaa gatgggtgtt 420
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<210> 22
 <211> 1145
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> Incyte ID No: 3201881CB1

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 tccctgatc agcgtaaccag ttgttgcctg tctgaacctc tgccagtcct ggagactgg 180
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 gctgcacagg cggaaaagtt ccagcaccc gggctgtaca tgcggcagga aaagccctcg 360
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 accag 1145

<210> 23
 <211> 3084
 <212> DNA
 <213> Homo sapiens

<220>
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 <223> Incyte ID No: 939000CB1

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 ttggctcgcc cggccaccat gaacaagaag aaaaaccgt tccttagggat gcccgcgc 180
 ctcggctacg tgccggggct gggccggggc gccactggct tcaccacgcg gtcagacatt 240
 gggcccgccc gtatgcacaa tgaccctgtg gatgatgcgc atgcacccccc aggcaagaga 300
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 <211> 3315
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> Incyte ID No: 2125677CB1

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 ttctcagaat tcagcgagaa gagagaggat gactgcgcga gaagaagcta gtttacgaac 180
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aggagacttc ctaaattatg ctctgtctct aatgcggctc cataatgatg agcattctga 300
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<210> 25

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<211> 1677

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

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<210> 26

<211> 997

<212> DNA

<213> Homo sapiens

<220>

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<210> 27

<211> 1481

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 3255641CB1

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<211> 303

<212> DNA

<213> Homo sapiens

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<220>
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 <213> Homo sapiens

<220>
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<220>
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 <213> Homo sapiens

<220>
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<220>
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 cattcctact aaggggctct gagttctgc ccccaagaatc attccaaccc acccaactgca 840
 aagactatga cagcatcaa tttcaggacc tgcagacagt acaggctaga taacccaccc 900
 aattttccca ctgtcctctg atccccctgt gacagaacct ttcagataa cgccctcacat 960
 cccaaagtcta tacccttacc tgaagaatgc tggttttcc tagccacett tctagcctcc 1020
 cacttgcctt gaaaggccaa gatcaagatg tcccccaaggc atcttgcatac cagcctgact 1080
 gctgctacat ctaatccccctt accaatgcct cctgtcccta aactccccag catactgatg 1140
 acagccctct ctgactttac cttgagatct gtcttcatac cttcccttc aaactaaca 1200
 aaacatttcc aataaaaata tcaagaatac 1229

<210> 35
 <211> 1455
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> Incyte ID No: 001273CB1

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<400> 35

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 ggcttcatt gcacaccgctg gctaatgccc cggccacggc tacagaaacg acctgccaag 120
 acgtcgccg gaccccccgtc ggcgggtacc cggcgattgt ggcctccatg acagccgaca 180
 gcaaagctgc acggctgcgg cggatcgagc gctggcaggc gacgggtgcac gctgcggagt 240
 cggtagacga gaagctgcga atccctacca agatcgagtt tatgaagttac atggtttacc 300
 cgcagaccc cgcgctgaat gccgaccgct ggtaccagta cttcaccaag accgtgttcc 360
 tgcgggtct gcccggcgc cccagcgagc cggagcccgaa gcccgaaccc gaacctgaac 420
 ctgcgtcggc cctcgccggc ctgcgtcggc tcgcctgcga ctgcctgctg caggagcact 480
 tctacctgcg ggcaggcggc cgcgtcggc acc gttacgagga gagcgaggc atatcttgc 540
 ccttccttggc tcagctggc tcaaccctcg tggcctcct cagccacac aaccggccc 600
 tggccgtgc cgcctcgat tataatgtcc cagttcattt ttactgggtg cgtggtaag 660
 aaattattcc tcgtggtcat cgaagaggc gaattgtga cttgcgatac cagatagatg 720
 ataaacccaa caaccagatt cgaatatcca agcaactcg agagttgtg ccattggatt 780
 attctgttcc tataatgtcc cccactataa aatgtaaacc agacaaactt ccattattca 840
 aacggcgtt tggaaaccac atatttggc gctaaaaac tgcagatcct tgctgttaag 900
 gtcacacccca gtttcatctg ttacctgaca aattaagaag gggaaaggc ttgagacaaa 960
 actgtgtcga tcagatagaa gttgttttta gagctaattgc tattgtcaagc ctttttgc 1020
 ggactggagc acaagctatg tatcaaggat tctggagtga agcagatgtt actcgaccc 1080
 ttgtctccca ggctgtgatc acagatggaa aataactttc ctttttgc taccagctaa 1140
 atactttggc actgactaca caagctgatc aaaataaccc tcgtaaaaat atatgttgg 1200
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 atgatgttct acttcagata gttcaatttc tactgtatag accaaaaagaa gaaaaatcac 1320
 agctgttggc aaactgaaaa agcatatgg attgagaact gtggaaatat ttaaatttta 1380
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 gacaaaaaaa aaaaaa 1455

<210> 36

<211> 2099

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 411831CB1

<400> 36

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 cccaagcgcc ggcggctcca gctccggac gacgaccacg acgacgacca cgacggggagg 180
 gatcctgatc ggctgtgatc tgcgttgcg agtttactt accatcgacc actctctgat 240
 tccggaggag aggctctcgcc ccacccatc catgcaggat gggctcgacc tgcccagtga 300
 gacggactta cgcatcctgg gctgcgttgg catccaggcc gccggcattc tcctccggct 360
 gccgcagggt gcgatggcaa cggggcagggt ttgtttcat cgtttttct actccaaatc 420
 ttctgtcaaa cacagtttcg agattgttgc tatggcttgcg attaatcttgcatcaaaaat 480
 cgaagaagca cctagaagaa taagagatgt gattaatgtt ttcaccacc tccgcccagg 540
 aagaggaaaa aggactccaa gccccctgtat cttgtatcag aactacatta acacaaaaaa 600
 tcaagttatc aaagcagaga ggagggtgtt aaaggagttt ggatgttgc ttcatgtcaa 660
 gcatcctcat aagatcattt ttatgtatcc acaagtcttta gaatgtgaac gtaatcaa 720
 cctgggttcaa actgccttggc attacatggc tgacagtctt cgaaccaatg tgtttgc 780
 atttcaacca gagactatag catgtgttgc catctacctt gcagcttagag cacttcagat 840
 tccgttgcgc actcgcccc attgggttct tcttttgcg actacagaag agggaaatcca 900
 gggaaatctgc atagaaaacac ttaggcttgc taccagaaaa aagccaaact atgaattact 960
 gggaaaaagaa gtagaaaaaaa gaaaagtagc cttacaagaa gccaaattaa aagccaaagg 1020

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attgaatccg gatggaactc cagcccttc aaccctgggt ggattttctc cagcctccaa 1080
 gccatcatca ccaagagaag taaaagctga agagaaatca ccaatctcca ttaatgtgaa 1140
 gacagtcaaa aaagaacctg aggatagaca acaggcttc aaaagccctt acaatggtgt 1200
 aagaaaagac agcaagagaa gtagaaatag cagaagtgcg agtcgatcga ggtcaagaac 1260
 acgatcacgt tctagatcac atactccaag aagacactat aataataggc ggagtcgatc 1320
 tggAACatac agctcgagat caagaagcag gtcccgact cacagtqaaa gccctcgaaag 1380
 acatcataat catgggtctc ctcaccttaa ggccaagcat accagagatg attaaaaaag 1440
 ttcaaacaga catggtcata aaaggaaaaa atctcggtct cgatctcaga gcaagtctcg 1500
 ggtactca gatgcagcca agaaacacag gcatgaaagg ggacatcata gggacaggcg 1560
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 agaactatgt taattttttt gcacattaaa atgccttagc agtataat taaaaaccat 1860
 ggtcaggttc aattgtactt tattatagtt gtgtattgtt tattgcata agaactggag 1920
 cgtgaattct gtaaaaatgt atcttatttt tatacagata aaattgcaga cactgttcta 1980
 tttaagtgg tattttttta aatgatggtg aatactttct taacactggt ttgtctgcat 2040
 gtgtaaagat ttttacaagg aaataaaaata caaatcttgc ttttctaaa aaaaaaaaaa 2099

<210> 37
 <211> 1363
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> Incyte ID No: 1520835CB1

<400> 37
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 cagggtgtct ggtggcaggc ctgtgcctat ccctgctgtc cccagggtgg gccccggggg 120
 tcaggagctc cagaagggcc agctggcat attctgagat tggccatcag cccccatttc 180
 tgctgcaaaac ctggtcagag ccagtgttcc ctccatgggg cctaaagaca gtgccaagt 240
 cctgcaccgt ggaccacagc cgagccactg ggcagccgt gatggtccc cgcaggagcg 300
 ctgtggaccg cgctctctgg gcagccctgt cctaggcctg gacacctgca gagcctggg 360
 ccacgtggat gggcagatcc tggccagct gcccggccctg acagaggagg aagaggagga 420
 gggcgcggg gccacctgt ccagggggcc tgccttcccc ggcattggct ctgaggagtt 480
 gctgtggcc tccttctatg actggccgct gactgcttag gtgccacccg agctgctggc 540
 tgctgcccgc ttcttccaca caggccatca ggacaagggt aggtgcttct tctgctatgg 600
 gggcctgcag agctgaaagc gccccggacg cccctggacg gagcatgcca agtggttccc 660
 cagctgttag ttctgtcc gtcataaaagg aagagacttt gtccacagtg tgcaggagac 720
 tcactcccag ctgctgggt cctgggaccc gtgggaagaa ccggaaagacg cagccccgt 780
 ggcggccctcc gtccctgcct ctgggtaccc tgagctgccc acacccagga gagagggtcca 840
 gtctgaaagt gcccaggagc caggagggtt cagtcagcc gagggccaga gggcgtgggt 900
 ggttcttgag ccccccaggag ccagggatgt ggaggcgcag ctgcggccgc tgcaggagga 960
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 ccgcagccgc gtgcgcaccc tcctgtccca ggccagggtgc catggccggc caggtgggt 1140
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 atggcagagc tgggttccat ccagcactga ccagccctga ttcccccacc accgcccagg 1260
 gtggagaagg aggcccttgc ttggcgtggg ggatggctt actgtacctg tttggatgct 1320
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<210> 38
 <211> 1465
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <223> Incyte ID No: 1902803CB1

<400> 38
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 tcagcatgga ggcaaaagga cagtggcaa gaccacgtcc ctccgaaggg agatggctgc 120
 ggggatgtat ttggaacatt atctggacag tattgaaaac cttcccttg aattacagag 180
 aaactttcag ctcatgaggg accttagacca aagaacagag gacctaagg ctgaaattga 240
 caagttggcc actgagtata ttagtagtgc ccgcagcctg agctccgagg aaaaattggc 300
 ccttctcaaa cagatccagg aagcctatgg caagtgcag gaatttggtg acgacaaggt 360
 gcagcttggc atgcagacct atgagatggt ggacaaacac attcggcggc tggacacaga 420
 cctggccgt tttgaggctg atctcaagga gaaacagatt gagtcaagtg actatgacag 480
 ctcttccagc aaaggcaaaa agaaaggccg gactcaaaag gagaagaaaag ctgctcgtgc 540
 tctttccaaa gggaaaaact cggatgaaga agcccccaag actgcccaga agaagttaaa 600
 gctcgtgcgc acaagtcctg agtatggat gcctcagtgc accttggca gtgtccaccc 660
 ctctgatgtg ttggatatgc ctgtggatcc caacgaaccc acctattgcc tttgtcacca 720
 ggtctctat ggagagatga ttggctgtga caaccctgtat tgttccattt agtggttcca 780
 ttttgcctgt gtggggctga caaccaagcc tcggggggaaa tggtttgcc cacgctgctc 840
 ccaagaacgg aagaagaaat agataaggcc ttggattcc aacacagttt cttccacatc 900
 ccctgacttg ggctagtggg cagaggaatg cctgtgctgg ggccagggggt tcagggagga 960
 gtggatggca cagtgcgtc atcccttctc ctccccctctc cccactcccc gtgctgaggc 1020
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 ggcatagcat gggggcagtc ccccagaccc ctccatccc ctcctgtgg tgagggtctag 1260
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<210> 39
 <211> 332
 <212> PRT
 <213> Mus musculus

<300>
 <308> GenBank ID No: g452276

<400> 39
 Met Ala Thr Pro Val Pro Pro Ser Pro Arg His Leu Arg Leu
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 Leu Arg Leu Leu Leu Ser Gly Leu Ile Leu Gly Ala Ala Leu Asn
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 Gly Ala Thr Ala Arg Arg Pro Asp Ala Thr Thr Cys Pro Gly Ser

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35	40	45
Leu Asp Cys Ala Leu Lys Arg Arg Ala Lys Cys Pro Pro Gly Ala		
50	55	60
His Ala Cys Gly Pro Cys Leu Gln Ser Phe Gln Glu Asp Gln Arg		
65	70	75
Gly Phe Cys Val Pro Arg Lys His Leu Ser Ser Gly Glu Gly Leu		
80	85	90
Pro Gln Pro Arg Leu Glu Glu Glu Ile Asp Ser Leu Ala Gln Glu		
95	100	105
Leu Ala Leu Lys Glu Lys Glu Ala Gly His Ser Arg Leu Thr Ala		
110	115	120
Gln Pro Leu Leu Glu Arg Ala Gln Lys Leu Leu Glu Pro Ala Ala		
125	130	135
Thr Leu Gly Phe Ser Gln Trp Gly Gln Arg Leu Glu Pro Gly Leu		
140	145	150
Pro Ser Thr His Gly Thr Ser Ser Pro Ile Pro His Thr Ser Leu		
155	160	165
Ser Ser Arg Ala Ser Ser Gly Pro Val Gln Met Ser Pro Leu Glu		
170	175	180
Pro Gln Gly Arg His Gly Asn Gly Leu Thr Leu Val Leu Ile Leu		
185	190	195
Ala Phe Cys Leu Ala Ser Ser Ala Ala Leu Ala Val Ala Ala Leu		
200	205	210
Cys Trp Cys Arg Leu Gln Arg Glu Ile Arg Leu Thr Gln Lys Ala		
215	220	225
Asp Tyr Ala Ala Thr Ala Lys Gly Pro Thr Ser Pro Ser Thr Pro		
230	235	240
Arg Ile Ser Pro Gly Asp Gln Arg Leu Ala His Ser Ala Glu Met		
245	250	255
Tyr His Tyr Gln His Gln Arg Gln Gln Met Leu Cys Leu Glu Arg		
260	265	270
His Lys Glu Pro Pro Lys Glu Leu Glu Ser Ala Ser Ser Asp Glu		
275	280	285
Glu Asn Glu Asp Gly Asp Phe Thr Val Tyr Glu Cys Pro Gly Leu		
290	295	300
Ala Pro Thr Gly Glu Met Glu Val Arg Asn Pro Leu Phe Asp His		
305	310	315
Ser Thr Leu Ser Ala Pro Val Pro Gly Pro His Ser Leu Pro Pro		
320	325	330
Leu Gln		

<210> 40
 <211> 268
 <212> PRT
 <213> Homo sapiens

<300>
 <308> GenBank ID No: g998357

<400> 40
 Met Ala Val Asn Val Tyr Ser Thr Ser Val Thr Ser Asp Asn Leu
 1 5 10 15

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Ser Arg His Asp Met Leu Ala Trp Ile Asn Glu Ser Leu Gln Leu
 20 25 30
 Asn Leu Thr Lys Ile Glu Gln Leu Cys Ser Gly Ala Ala Tyr Cys
 35 40 45
 Gln Phe Met Asp Met Leu Phe Pro Gly Ser Ile Ala Leu Lys Lys
 50 55 60
 Val Lys Phe Gln Ala Lys Leu Glu His Glu Tyr Ile Gln Asn Phe
 65 70 75
 Lys Ile Leu Gln Ala Gly Phe Lys Arg Met Gly Val Asp Lys Ile
 80 85 90
 Ile Pro Val Asp Lys Leu Val Lys Gly Lys Phe Gln Asp Asn Phe
 95 100 105
 Glu Phe Val Gln Trp Phe Lys Lys Phe Phe Asp Ala Asn Tyr Asp
 110 115 120
 Gly Lys Asp Tyr Asp Pro Val Ala Ala Arg Gln Gly Gln Glu Thr
 125 130 135
 Ala Val Ala Pro Ser Leu Val Ala Pro Ala Leu Asn Lys Pro Lys
 140 145 150
 Lys Pro Leu Thr Ser Ser Ala Ala Pro Gln Arg Pro Ile Ser
 155 160 165
 Thr Gln Arg Thr Ala Ala Ala Pro Lys Ala Gly Pro Gly Val Val
 170 175 180
 Arg Lys Asn Pro Gly Val Gly Asn Gly Asp Asp Glu Ala Ala Glu
 185 190 195
 Leu Met Gln Gln Val Asn Val Leu Lys Leu Thr Val Glu Asp Leu
 200 205 210
 Glu Lys Glu Arg Asp Phe Tyr Phe Gly Lys Leu Arg Asn Ile Glu
 215 220 225
 Leu Ile Cys Gln Glu Asn Glu Gly Glu Asn Asp Pro Val Leu Gln
 230 235 240
 Arg Ile Val Asp Ile Leu Tyr Ala Thr Asp Glu Gly Phe Val Ile
 245 250 255
 Pro Asp Glu Gly Gly Pro Gln Glu Glu Gln Glu Glu Tyr
 260 265

<210> 41
 <211> 418
 <212> PRT
 <213> Mus musculus

<300>
 <308> GenBank ID No: g455719

<400> 41

Met	Gly	Glu	Asp	Ala	Ala	Gln	Ala	Glu	Lys	Phe	Gln	His	Pro	Asn
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Thr	Asp	Met	Leu	Gln	Glu	Lys	Pro	Ser	Ser	Pro	Ser	Pro	Met	Pro
	20							25				30		
Ser	Ser	Thr	Pro	Ser	Pro	Ser	Leu	Asn	Leu	Gly	Ser	Thr	Glu	Glu
	35							40				45		
Ala	Ile	Arg	Asp	Asn	Ser	Gln	Val	Asn	Ala	Val	Thr	Val	His	Thr
	50							55				60		

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Leu	Leu	Asp	Lys	Leu	Val	Asn	Met	Leu	Asp	Ala	Val	Arg	Glu	Asn
				65					70					75
Gln	His	Asn	Met	Glu	Gln	Arg	Gln	Ile	Asn	Leu	Glu	Gly	Ser	Val
				80					85					90
Lys	Gly	Ile	Gln	Asn	Asp	Leu	Thr	Lys	Leu	Ser	Lys	Tyr	Gln	Ala
				95					100					105
Ser	Thr	Ser	Asn	Thr	Val	Ser	Lys	Leu	Leu	Glu	Lys	Ser	Arg	Lys
				110					115					120
Val	Ser	Ala	His	Thr	Arg	Ala	Val	Arg	Glu	Arg	Leu	Glu	Arg	Gln
				125					130					135
Cys	Val	Gln	Val	Lys	Arg	Leu	Glu	Asn	Asn	His	Ala	Gln	Leu	Leu
				140					145					150
Arg	Arg	Asn	His	Phe	Lys	Val	Leu	Ile	Phe	Gln	Glu	Glu	Ser	Glu
				155					160					165
Ile	Pro	Ala	Ser	Val	Phe	Val	Lys	Glu	Pro	Val	Pro	Ser	Ala	Ala
				170					175					180
Glu	Gly	Lys	Glu	Glu	Leu	Ala	Asp	Glu	Asn	Lys	Ser	Leu	Glu	Glu
				185					190					195
Thr	Leu	His	Asn	Val	Asp	Leu	Ser	Ser	Asp	Asp	Glu	Leu	Pro	Arg
				200					205					210
Asp	Glu	Glu	Ala	Leu	Glu	Asp	Ser	Ala	Glu	Glu	Lys	Met	Glu	Glu
				215					220					225
Ser	Arg	Ala	Glu	Lys	Ile	Lys	Arg	Ser	Ser	Leu	Lys	Lys	Val	Asp
				230					235					240
Ser	Leu	Lys	Lys	Ala	Phe	Ser	Arg	Gln	Asn	Ile	Glu	Lys	Lys	Met
				245					250					255
Asn	Lys	Leu	Gly	Thr	Lys	Ile	Val	Ser	Val	Glu	Arg	Arg	Glu	Lys
				260					265					270
Ile	Lys	Lys	Ser	Leu	Thr	Pro	Asn	His	Gln	Lys	Ala	Ser	Ser	Gly
				275					280					285
Lys	Ser	Ser	Pro	Phe	Lys	Val	Ser	Pro	Leu	Ser	Phe	Gly	Arg	Lys
				290					295					300
Lys	Val	Arg	Glu	Gly	Glu	Ser	Ser	Val	Glu	Asn	Glu	Thr	Lys	Leu
				305					310					315
Glu	Asp	Gln	Met	Gln	Glu	Asp	Arg	Glu	Glu	Gly	Ser	Phe	Thr	Glu
				320					325					330
Gly	Leu	Ser	Glu	Ala	Ser	Leu	Pro	Ser	Gly	Leu	Met	Glu	Gly	Ser
				335					340					345
Ala	Glu	Asp	Ala	Glu	Lys	Ser	Ala	Arg	Arg	Gly	Asn	Asn	Ser	Ala
				350					355					360
Val	Gly	Ser	Asn	Ala	Asp	Leu	Thr	Ile	Glu	Glu	Asp	Glu	Glu	Glu
				365					370					375
Glu	Pro	Val	Ala	Leu	Gln	Gln	Ala	Gln	Gln	Val	Arg	Tyr	Glu	Ser
				380					385					390
Gly	Tyr	Met	Leu	Asn	Ser	Glu	Glu	Met	Glu	Glu	Pro	Ser	Glu	Lys
				395					400					405
Gln	Val	Gln	Pro	Ala	Val	Leu	His	Val	Asp	Gln	Thr	Ala		
				410					415					

<210> 42
 <211> 142
 <212> PRT
 <213> Homo sapiens

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<300>

<308> GenBank ID No: g2565275

<400> 42

Met	Ser	Tyr	Met	Leu	Pro	His	Leu	His	Asn	Gly	Trp	Gln	Val	Asp
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Gln	Ala	Ile	Leu	Ser	Glu	Glu	Asp	Arg	Val	Val	Val	Ile	Arg	Phe
					20				25					30
Gly	His	Asp	Trp	Asp	Pro	Thr	Cys	Met	Lys	Met	Asp	Glu	Val	Leu
					35				40					45
Tyr	Ser	Ile	Ala	Glu	Lys	Val	Lys	Asn	Phe	Ala	Val	Ile	Tyr	Leu
					50				55					60
Val	Asp	Ile	Thr	Glu	Val	Pro	Asp	Phe	Asn	Lys	Met	Tyr	Glu	Leu
					65				70					75
Tyr	Asp	Pro	Cys	Thr	Val	Met	Phe	Phe	Arg	Asn	Lys	His	Ile	
					80				85					90
Met	Ile	Asp	Leu	Gly	Thr	Gly	Asn	Asn	Asn	Lys	Ile	Asn	Trp	Ala
					95				100					105
Met	Glu	Asp	Lys	Gln	Glu	Met	Val	Asp	Ile	Ile	Glu	Thr	Val	Tyr
					110				115					120
Arg	Gly	Ala	Arg	Lys	Gly	Arg	Gly	Leu	Val	Val	Ser	Pro	Lys	Asp
					125				130					135
Tyr	Ser	Thr	Lys	Tyr	Arg	Tyr								
				140										

<210> 43

<211> 464

<212> PRT

<213> Drosophila melanogaster

<300>

<308> GenBank ID NO: g3688609

<400> 43

Met	Glu	Ala	Asp	Gly	Leu	Thr	Asn	Glu	Gln	Thr	Glu	Lys	Val	Leu
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Gln	Phe	Gln	Asp	Leu	Thr	Gly	Ile	Glu	Asp	Met	Asn	Val	Cys	Arg
					20				25					30
Asp	Val	Leu	Ile	Arg	His	Gln	Trp	Asp	Leu	Glu	Val	Ala	Phe	Gln
					35				40					45
Glu	Gln	Leu	Asn	Ile	Arg	Glu	Gly	Arg	Pro	Thr	Met	Phe	Ala	Ala
					50				55					60
Ser	Thr	Asp	Val	Arg	Ala	Pro	Ala	Val	Leu	Asn	Asp	Arg	Phe	Leu
					65				70					75
Gln	Gln	Val	Phe	Ser	Ala	Asn	Met	Pro	Gly	Gly	Arg	Thr	Val	Ser
					80				85					90
Arg	Val	Pro	Ser	Gly	Pro	Val	Pro	Arg	Ser	Phe	Thr	Gly	Ile	Ile
					95				100					105
Gly	Tyr	Val	Ile	Asn	Phe	Val	Phe	Gln	Tyr	Phe	Tyr	Ser	Thr	Leu
					110				115					120
Thr	Ser	Ile	Val	Ser	Ala	Phe	Val	Asn	Leu	Gly	Gly	Gly	Asn	Glu
					125				130					135

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Ala Arg Leu Val Thr Asp Pro Leu Gly Asp Val Met Lys Phe Ile
 140 145 150
 Arg Glu Tyr Tyr Glu Arg Tyr Pro Glu His Pro Val Phe Tyr Gln
 155 160 165
 Gly Thr Tyr Ala Gln Ala Leu Asn Asp Ala Lys Gln Glu Leu Arg
 170 175 180
 Phe Leu Ile Val Tyr Leu His Lys Asp Pro Ala Lys Asn Pro Asp
 185 190 195
 Val Glu Ser Phe Cys Arg Asn Thr Leu Ser Ala Arg Ser Val Ile
 200 205 210
 Asp Tyr Ile Asn Thr His Thr Leu Leu Trp Gly Cys Asp Val Ala
 215 220 225
 Thr Pro Glu Gly Tyr Arg Val Met Gln Ser Ile Thr Val Arg Ser
 230 235 240
 Tyr Pro Thr Met Val Met Ile Ser Leu Arg Ala Asn Arg Met Met
 245 250 255
 Ile Val Gly Arg Phe Glu Gly Asp Cys Thr Pro Glu Glu Leu Leu
 260 265 270
 Arg Arg Leu Gln Ser Val Thr Asn Ala Asn Glu Val Trp Leu Ser
 275 280 285
 Gln Ala Arg Ala Asp Arg Leu Glu Arg Asn Phe Thr Gln Thr Leu
 290 295 300
 Arg Arg Gln Gln Asp Glu Ala Tyr Glu Gln Ser Leu Leu Ala Asp
 305 310 315
 Glu Glu Lys Glu Arg Gln Arg Gln Arg Glu Arg Asp Ala Val Arg
 320 325 330
 Gln Ala Glu Glu Ala Val Glu Gln Ala Arg Arg Asp Val Glu Leu
 335 340 345
 Arg Lys Glu Glu Ile Ala Arg Gln Lys Ile Glu Leu Ala Thr Leu
 350 355 360

 Val Pro Ser Glu Pro Ala Ala Asp Ala Val Gly Ala Ile Ala Val
 365 370 375
 Val Phe Lys Leu Pro Ser Gly Thr Arg Leu Glu Arg Arg Phe Asn
 380 385 390
 Gln Thr Asp Ser Val Leu Asp Val Tyr His Tyr Leu Phe Cys His
 395 400 405
 Pro Asp Ser Pro Asp Glu Phe Glu Ile Thr Thr Asn Phe Pro Lys
 410 415 420
 Arg Val Leu Phe Ser Lys Ala Asn Leu Asp Ala Ala Gly Glu Thr
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20 25 30
Ala Pro Pro Pro Gln Arg Pro Phe Arg Val Cys Asp His Lys Arg
35 40 45
Thr Val Arg Lys Gly Leu Thr Ala Ala Ser Leu Gln Glu Leu Leu
50 55 60
Asp Lys Val Leu Glu Thr Leu Leu Arg Gly Val Leu Thr Leu
65 70 75
Val Leu Glu Glu Asp Gly Thr Ala Val Asp Ser Glu Asp Phe Phe
80 85 90
Gln Leu Leu Glu Asp Asp Thr Cys Leu Met Val Leu Glu Gln Gly
95 100 105
Gln Ser Trp Ser Pro Lys Ser Gly Met Leu Ser Tyr Gly Leu Gly
110 115 120
Arg Glu Lys Pro Lys His Ser Lys Asp Ile Ala Arg Ile Thr Phe
125 130 135
Asp Val Tyr Lys Gln Asn Pro Arg Asp Leu Phe Gly Ser Leu Asn
140 145 150
Val Lys Ala Thr Phe Tyr Gly Leu Tyr Ser Met Ser Cys Asp Phe
155 160 165
Gln Gly Val Gly Pro Lys Arg Val Leu Arg Glu Leu Leu Arg Gly
170 175 180
Thr Ser Ser Gln Leu Gln Gly Leu Gly His Met Leu Leu Gly Ile
185 190 195
Ser Ser Thr Leu Arg His Val Val Glu Gly Ala Asp Arg Trp Gln
200 205 210
Trp His Gly Gln Arg His Leu His Ser
215